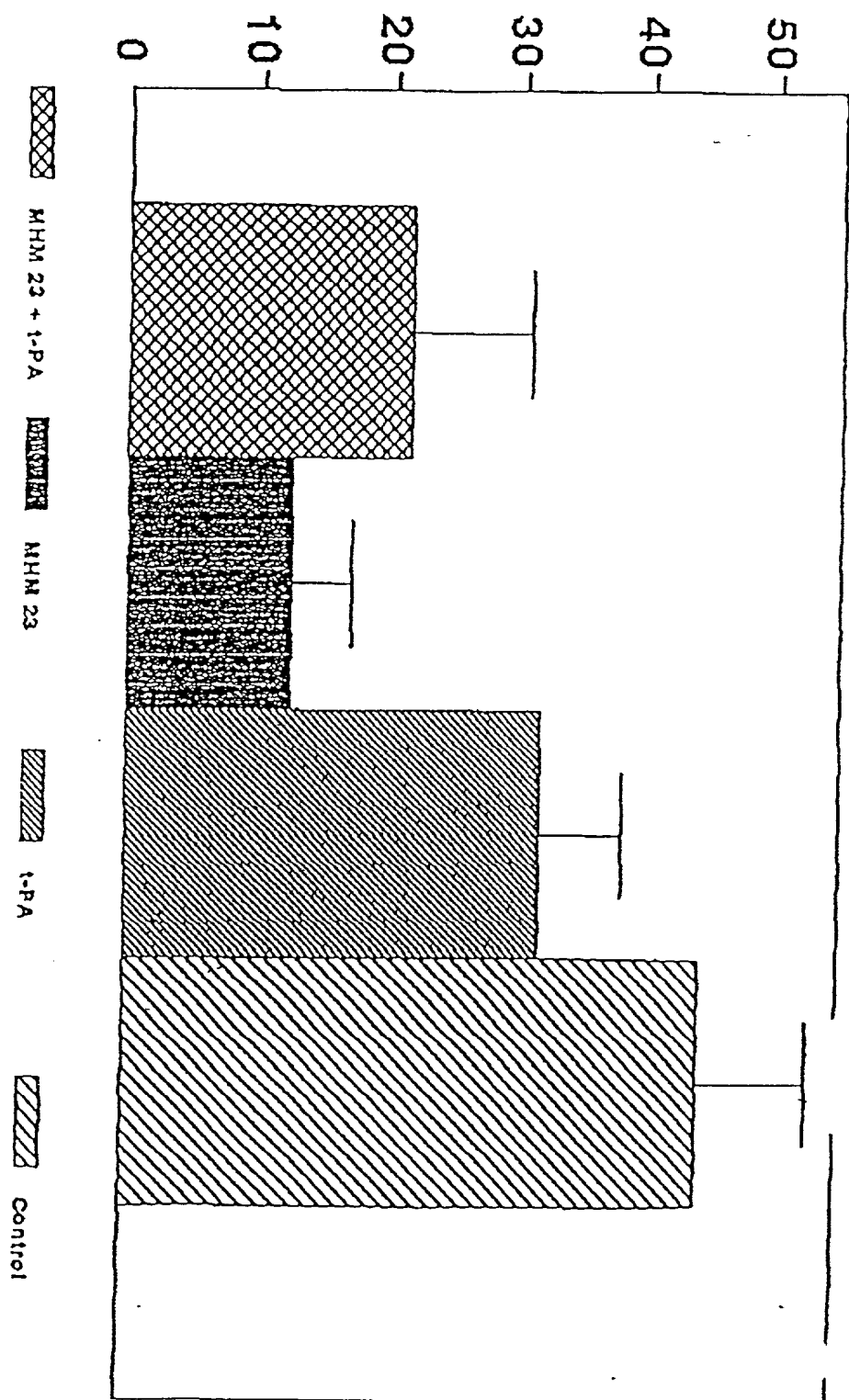
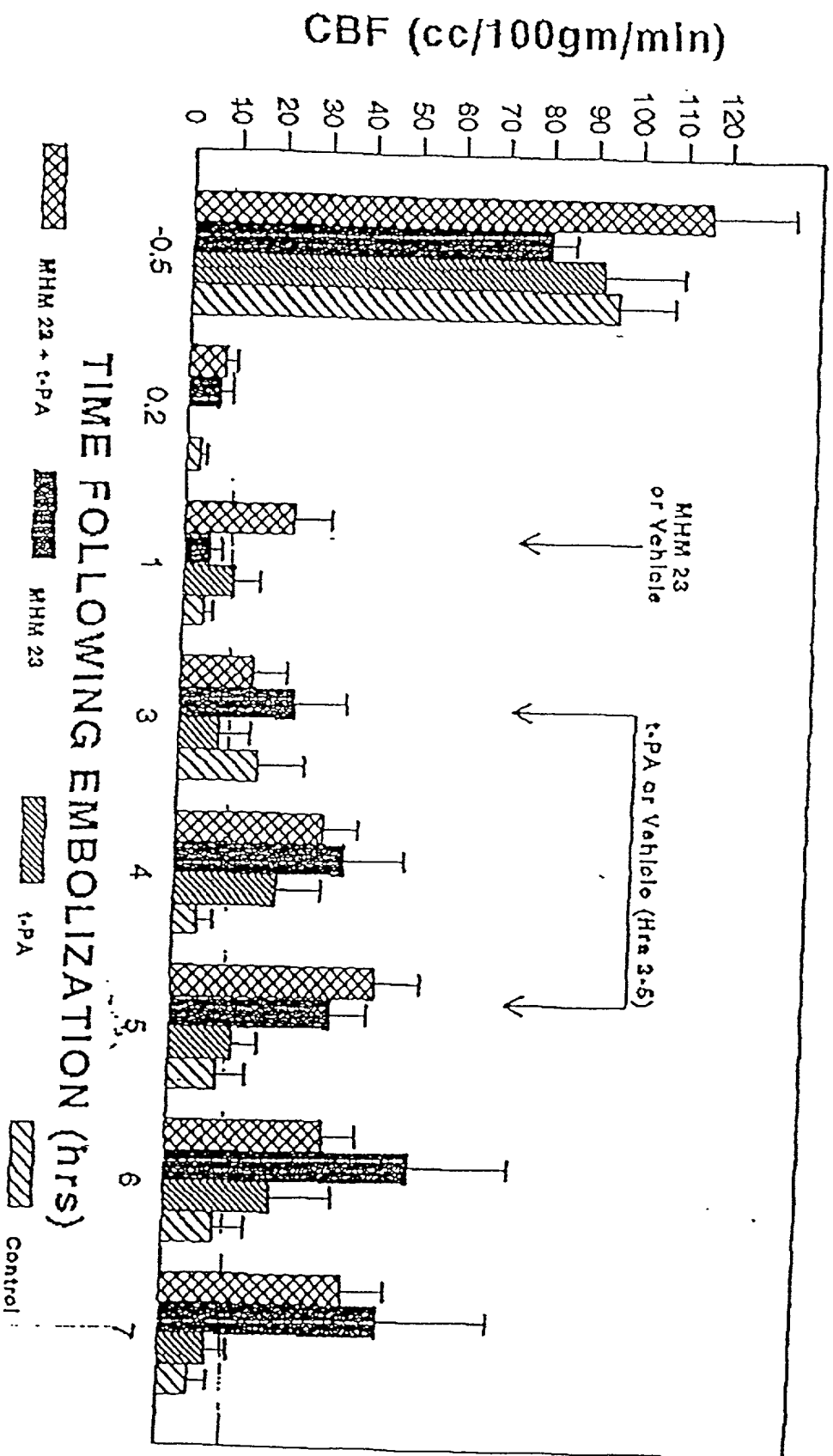


Infarct Size (% Hemisphere)



0000442244220000






卷一百一十五

KHM 23



PA



Control

TIME FOLLOWING EMBOLIZATION (hrs)

MMH 23
or Vehicle

1-PA or Vehicle (Hrs 3-5)

hIgG₁ Sequence of human IgG1 CH1 domain
hIgG₂ Sequence of human IgG2 CH1 domain
hIgG₃ Sequence of human IgG3 CH1 domain
hIgG₄ Sequence of human IgG4 CH1 domain
humκ Sequence of human kappa CL domain
humλ Sequence of human lambda CL domain

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              114                      128                      139
              |                      |                      |
hIgG1      A S T K G P S V F P L A P S S K S T S G G T A A L
hIgG2      A S T K G P S V F P L A P C S R S T S E S T A A L
hIgG3      A S T K G P S V F P L A P C S R S T S G G T A A L
hIgG4      A S T K G P S V F P L A P C S R S T S E S T A A L

              108                      122                      131
              |                      |                      |
humκ       R T V A A P S V F I F P P S D E Q L K S G T A S V
humλ       Q P K A A P S V T L F P P S S E E L Q A N K A T L
  
```

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Fabv1b      A S T K G P S V F P L A P S P K N S S M I S N T P A L
of interest                                P K N S S M I S N T P
  
```

most important *** *

```

hIgG1      G C L V K D Y F P E P V T V S W N S G A L T S
hIgG2      G C L V K D Y F P E P V T V S W N S G A L T S
hIgG3      G C L V K D Y F P E P V T V S W N S G A L T S
hIgG4      G C L V K D Y F P E P V T V S W N S G A L T S

humκ       V C L L N N F Y P R E A K V Q W K V D N A L Q
humλ       V C L I S D F Y P G A V T V A W K A D S S P V
  
```

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Fabv1b      G C L V K D Y F P E P V T V S W N S G A L T S

hIgG1      G V H T F P A V L Q S S G - - - L Y S L S S V
hIgG2      G V H T F P A V L Q S S G - - - L Y S L S S V
  
```

Figure 4A

hIgG ₃	G V H T F P A V L Q S S G - - - L Y S L S S V
hIgG ₄	G V H T F P A V L Q S S G - - - L Y S L S S V
humκ	S G N S Q E S V T E Q D S K D S T Y S L S S T
humλ	K A G V E T T T P S K Q S N N - K Y A A S S Y

Fabv1b	G V H T F P A V L Q S S G - - - L Y S L S S V

	193	200	203	
	I	I	I	
hIgG ₁	V T V P S S S L G T - Q T Y I C N V N H K P S			
hIgG ₂	V T V P S S N F G T - Q T Y T C N V D H K P S			
hIgG ₃	V T V P S S S L G T - Q T Y T C N V N H K P S			
hIgG ₄	V T V P S S S L G T - K T Y T C N V D H K P S			
	181		190	
	I		I	
humκ	L T L S K A D Y E K H K V Y A C E V T H Q G L			
humλ	L S L T P E Q W K S H R S Y S C Q V T H E G S			

Fabv1b	V T V P H Q S L G T - Q T Y I C N V N H K P S			
of interest	H Q N L S D G K			
most important	* * *		* *	

hIgG ₁	N T K V D K R V - - -
hIgG ₂	N T K V D K T V - - -
hIgG ₃	N T K V D K R V - - -
hIgG ₄	N T K V D K R V - - -
humκ	S S P V T K S F N R G E C
humλ	T V E K T V A P T E C S

Fabv1b	N T K V D K R V - - -

Figure 4B